

Table S1. *S. cerevisiae* strains used in this study

Name	Relevant Genotype	Source
RSY1533	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15</i>	Euroscarf
RSY2385	<i>Mat a leu2Δ0 ura3Δ0 his3Δ0 faa1::KanMX faa4::KanMX</i>	This study
RSY2965	<i>Mataα leu2 lys2 ura3 ole1^{ts} end3::KanMX</i>	This study
RSY2975	<i>Mat a leu2 lys2 ura3 ole1^{ts} vrp1::KanMX</i>	This study
RSY2976	<i>Mat a leu2 lys2 ura3 ole1^{ts} srv2::KanMX</i>	This study
RSY2997	<i>Mat a leu2 lys2 ura3 ole1^{ts} ypk1::KanMX</i>	This study
RSY3088	<i>Mat a leu2 lys2 ura3 ole1^{ts} faa1::KanMX faa4::KanMX</i>	This study
RSY3090	<i>Mataα leu2 lys2 ura3 ole1^{ts}</i>	This study
RSY3108	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15 ypk1::KanMX FAA4-GFP-HIS3</i>	This study
RSY3109	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15 FAA1-GFP-HIS3</i>	This study
RSY3110	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15 FAA4-GFP-HIS3</i>	This study
RSY3112	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15 ypk1::KanMX FAA1-GFP-HIS3</i>	This study
RSY3524 (LHY291)	<i>Mata his3 leu2 lys2 trp1 ura3 bar1-1</i>	deHart et al, 2002
RSY3526 (LHY2563)	<i>Mata ypk1Δ::KanMX his3::YPK1::HIS3 GAL2 his3 leu2 lys2 trp1 ura3 bar1-1</i>	deHart et al, 2002
RSY3527 (LHY 2564)	<i>Mata ypk1::KanMX his3::YPK1^{K376R}::HIS3 leu2 met15 ura3 bar1-1</i>	deHart et al, 2002
RSY3939	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15 ypk1::KanMX YEp351-ADH-SGK</i>	This study
RSY3948	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15 ypk1::KanMX pGREG506-Erg6-RFP</i>	This study
RSY4375	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15 ypk1::KanMX pGREG506-GAL-SGK1</i>	This study
RSY4379	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15 ypk1::KanMX</i>	Euroscarf
RSY4462	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15 end3::KanMX</i>	Euroscarf
RSY4463	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15 srv2::KanMX</i>	Euroscarf
RSY4464	<i>Mataα leu2Δ0 ura3Δ0 his3Δ0 lys2Δ0 MET15 vrp1::KanMX</i>	Euroscarf

Fig. S1. Ypk1 does not affect the localization of Faa1 and Faa4. The subcellular localization of GFP-tagged versions of Faa1 and Faa4 was examined by fluorescence microscopy in wild-type and *ypk1Δ* mutant cells. Scale bar: 5 μ m.

